

Title (en)
ANALYSIS OF FRAGMENTATION PATTERNS OF CELL-FREE DNA

Title (de)
ANALYSE VON FRAGMENTIERUNGSMUSTERN VON ZELLFREIER DNA

Title (fr)
ANALYSE DE MODÈLES DE FRAGMENTATION D'ADN ACELLULAIRE

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Application
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Abstract (en)

A method of analyzing a biological sample, comprising: identifying at least one genomic region having a fragmentation pattern specific to a first tissue type; analyzing a plurality of cell-free DNA molecules from the biological sample, the biological sample including cell free DNA molecules from a plurality of tissues types that includes the first tissue type, wherein analyzing a cell-free DNA molecule includes: determining a genomic position in a reference genome corresponding to at least one end of the cell-free DNA molecule; identifying a first set of first genomic positions, each first genomic position having a local minimum of ends of cell-free DNA molecules corresponding to the first genomic position; identifying a second set of second genomic positions, each second genomic position having a local maximum of ends of cell-free DNA molecules corresponding to the second genomic position; determining a first number of cell-free DNA molecules ending on any one of the first genomic positions in any one of the at least one genomic region; determining a second number of cell-free DNA molecules ending on any one of the second genomic positions in any one of the at least one genomic region; computing a separation value using the first number and the second number; and determining a classification of a proportional contribution of the first tissue type by comparing the separation value to one or more calibration values determined from one or more calibration samples whose proportional contributions of the first tissue type are known.

IPC 8 full level
C12Q 1/6886 (2018.01)

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